

09069-070501

CCGGGGGAGGCTCTTAGGATGTTGTGCTCCGCGGGGCTCAGACGAAATCTTCTGTGAATGGAAG
AAATGCTTCCAAGCAAACAGCCACTACCAGAACAACCTGAGAAAGAGGCCAGAGCGCGAGTTCTC
AAACCCTGATTTTCGCAGGAGCCGGAGGGGGGATATTGGAGAGAAGGTATTTCCAGTCACGCGCAG
TAACAGACCAGCCAAGGACCAGGACTGGAGTTCTGTTCTACAACGGTGGAACAGTGAACGGTCT
CCAAAGAGATGGAGTACGACGCTTACAACGACTCCGGCATCTATGATGATGAGTACTCTGATGG
CTTTGGCTACTTTGTGGACTTGGAGGAGGCGAGTCCGTGGGAGGCCAAGGTGGCCCCGGTCTTC
CTGGTGGTGATCTACAGCTTGGTGTGCTTCCTCGGTCTCCTAGGCAACGGCCTGGTGATTGTCA
TCGCCACCTTCAAGATGAAGAAGACCGTGAACACTGTGTGGTTTGTCAACCTGGCTGTGGCCGA
CTTCCTGTTCAACATCTTTTTGCCGATGCACATCACCTACGCGGCCATGGACTACCACTGGGTG
TTCGGGAAGGCCATGTGCAAGATCAGCAACTTCTTGCTCAGCCACAACATGTACACCAGCGTCT
TCCTGCTGACTGTCATCAGCTTTGACCGCTGCATCTCCGTGCTGCTCCCCGTCTGGTCCCAGAA
CCACCGCAGCATCCGCCTGGCCTACATGACCTGCTCGGCCGTCTGGGTCTGGCTTTCTTCTTG
AGCTCCCCGTCCCTTGCTCTCCGGGACACCGCCAACATTCATGGGAAGATAACCTGCTTCAACA
ACTTCAGCTTGGCCGCGCCTGAGTCCTCCCCACATCCCGCCCACTCGCAAGTAGTTTCCACAGG
GTACAGCAGACACGTGGCGGTCACCTGTCACCCGCTTCCTTTGCGGCTTCCTGATCCCCGTCTTC
ATCATCACGGCCTGCTACCTTACCATCGTCTTCAAGCTGCAGCGCAACCGCCTGGCCAAGAACA
AGAAGCCCTTCAAGATCATCATCACCATCATCATCACCTTCTTCTCTGCTGGTGCCCCTACCA
CACCCTCTACCTGCTGGAGCTCCACCACACAGCTGTGCCAAGCTCTGTCTTCAGCCTGGGGCTA
CCCCTGGCCACGGCCGTCGCCATCGCCAACAGCTGCATGAACCCCATTTCTGTACGTCTTCATGG
GCCACGACTTCAGAAAATTCAGGTGGCCCTCTTCTCCCGCCTGGCCAACGCCCTGAGTGAGGA
CACAGGCCCCCTCCTCCTACCCCAGTCACAGGAGCTTCACCAAGATGTCGTCTTTGAATGAGAAG
GCTTCGGTGAATGAGAAGGAGACCAGTACCCTCTGAACCTCACCTGGGAATGTCCCCCAAAGGT
GCCACGGCCCAGGGACGCCTAGGGACTTGTCTCCGGAAGTGGGAGACATGCCGGGAGCCTTTGG
GAATGCTCCAACGCCCCACTGAATTTTGCACAAGGCGGCTCATGTTTAAAGTGGGGTTCCCAAGT
GTGGACACTCTTCCAGTAAAATGGCAGGCAAGCAACCCGAGCTTCTACAACAGGAGCAGGGGAC
CGACTGTGACTGACTCAGAAAAGGGAGCATTTCTGAAGCCAAGACTTGAGCTGTGACCAACATA
CAGGCCAACATACAGATGTCGCCGTGCATGCCCTGAACATGCTGCGCAGTTTTCTGTGGGTGAG
GAAGTTACCGCAAACCCATTGCAGACCTGTTATGGCAACATGACAGTCAAACCAACAAAGCCCA
ATACACCCCAACATCCTCCAAGACCTTGACTTTGGATTTCAGAAGAACGGGGGGTGGGGGGAAC
GAGGACCTGAGGGTTAATTTTCGAGCTTGGCGAAGCC (SEQ ID NO:1)

FIGURE 1

underlined = deleted in targeting construct

[] = sequence flanking Neo insert in targeting construct

CCGGGGGAGGCTCTTAGGATGTTGTGCTCCGCGGGGCTCAGACGAAATCTTCTGTGAATG
GAAGAAATGCTTCCAAGCAAACAGCCACTACCAGAACAACCTGAGAAAGAGGCCAGAGCGC
GAGTTCTCAAACCCCTGATTTTCGCAGGAGCCGGAGGGGGATATTGGAGAGAAGGTATTTCC
AGTCACGCGCAGTAACAGACCAGCCAAGGACCAGGACTGGAGTTCTGTTCTACAACGGTG
GAACAGTGAACGGTCTCCAAAG [AGATGGAGTACGACGCTTACAACGACTCCGGCATCTA
TGATGATGAGTACTCTGATGGCTTTGGCTACTTTGTGGACTTGGAGGAGGCGAGTCCGTG
GGAGGCCAAGGTGGCCCCGGTCTTCCTGGTGGTGATCTACAGCTTGGTGTGCTTCCTCGG
TC] TCCTAGGCAACGGCCTGGTGATTGTCATCGCCACCTTCAAGATGAAGAAGACCGTGA
ACACTGTGTGGTTTGTCAACCTGGCTGTGGCCGACTTCCTGTTCAACATCTTTTTGCCGA
TGCACATCACCTACGCGGCCATGGACTACCCTGGGTG [TTCGGGAAGGCCATGTGCAAG
ATCAGCAACTTCTTGCTCAGCCACAACATGTACACCAGCGTCTTCCTGCTGACTGTCATC
AGCTTTGACCGCTGCATCTCCGTGCTGCTCCCCGTCTGGTCCCAGAACCACCGCAGCATC
CGCCTGGCCTACATGACCTGCTCGGCCGTCTGGGTCCTGGCTTTCTTCTTGAGCTCCCCG
TCCCTTGCTTCCGGGACACCGCCAACATTCATGGGAAGATAACCTGCTTCAACAACCTC
AGCTTGGCCGCGCCTGAGTCTTCCCCACATCCCCGCCACTCGCAAGTAGTTTCCACAGGG
TACAGCAGACACGTGGCGGTCACTGTCACCCGCTTCCTTTGCGGCTTCCTGATCCCCGTC
TTCATCATCACGGCCTGCTACCTTACCATCGTCTTCAAGCTGCAGCGCAACCGCCTGGCC
AAGAACAAGAAGCCCTTCAAGATCATCATCACCATCATCATCACCTTCTTCTCTGCTGG
TGCCCCCTACCACACCCTCTACCTGCTGGAGCTCCACCACACAGCTGTGCCAAGCTCTGTC
TTCAGCCTGGGGCTACCCCTGGCCACGGCCGTGCGCATCGCCAACAGCTGCATGAACCCC
ATTCTGTACGTCTTCATGGGCCACGACTTCAGAAAATTCAAGGTGGCCCTCTTCTCCCCG
CTGGCCAACGCCCTGAGTGAGGACACAGGCCC] CTCCTCCTACCCAGTCACAGGAGCTT
CACCAAGATGTCGTCTTTGAATGAGAAGGCTTCGGTGAATGAGAAGGAGACCAGTACCCT
CTGAACCTCACCTGGGAATGTCCCCCAAAGGTGCCACGGCCCAGGGACGCCTAGGGACTT
GTCTCCGGAAGTGGGAGACATGCCGGGAGCCTTTGGGAATGCTCCAACGCCCACTGAATT
TTGCACAAGGCGGCTCATGTTTTAAGTGGGGTTCCCAAGTGTGGACACTCTTCCAGTAAA
ATGGCAGGCAAGCAACCCGAGCTTCTACAACAGGAGCAGGGGACCGACTGTGACTGACTC
AGAAAAGGGAGCATTTCTGAAGCCAAGACTTGAGCTGTGACCAACATACAGGCCAACATA
CACGATGTCGCCGTGCATGCCCTGAACATGCTGCGCAGTTTTTCGTGGGTGAGGAAGTTAC
CGCAAACCCATTGCAGACCTGTTATGGCAACATGACAGTCAAACCAACAAAGCCCAATAC
ACCCCAACATCCTCCAAGACCTTGACTTTGGATTTCAGAAGAACGGGGGGTGGGGGAAC
GAGGACCTGAGGGTTAATTTTCGAGCTTGCGGAAGCC

FIGURE 2A

Gene Sequence Structure
*

422 bp

Sequence Deleted

576 bp

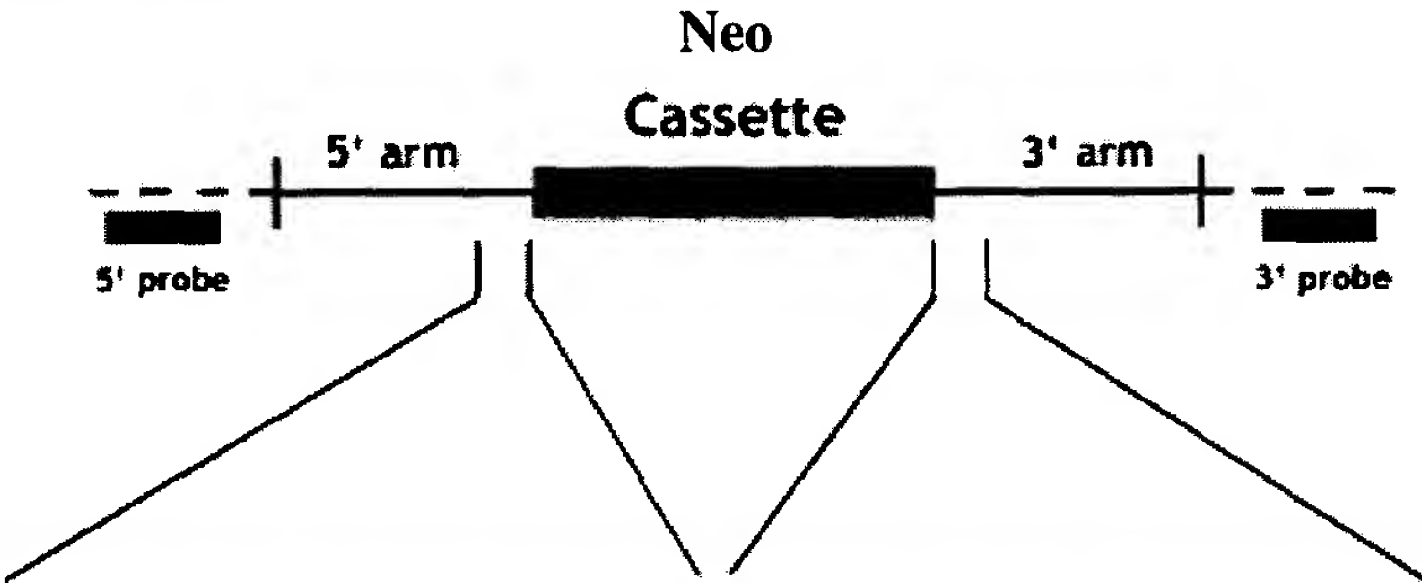
Size of full-length
cDNA: 1892 bp



Targeting Vector* (genomic sequence)

Construct Number: 993

Arm Length:
5': 2.3 kb
3': 1.9 kb



<p>5' >CCACAGAGGTCCTCAGCCTGT GACCCTGTCTTCCCTCACAGAGAT GGAGTACGACGCTTACAACGACTC CGGCATCTATGATGATGAGTACTC TGATGGCTTTGGCTACTTTGTGGA CTTGGAGGAGGCGAGTCCGTGGA GGCCAAGGTGGCCCCGGTCTTCCT GGTGGTGATCTACAGCTTGGTGTG CTTCCTCGGTC<3' (SEQ ID NO:2)</p>	<p>5' >TTCGGGAAGGCCATGTGCAAG ATCAGCAACTTCTTGCTCAGCCAC AACATGTACACCAGCGTCTTCCTG CTGACTGTCATCAGCTTTGACCGC TGCATCTCCGTGCTGCTCCCCGTC TGGTCCCAGAACCACCGCAGCATC CGCCTGGCCTACATGACCTGCTCG GCCGTCTGGGTCCTGGCTTTCTTC TTGAGCTCCCC<3' (SEQ ID NO:3)</p>
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Targeting Vector
Endogenous Locus

* Not drawn to scale

FIGURE 2B